INFORMATION DISCLOSURE STATEMENT BY APPLICANT

Application No. May 3, 2002 Filing Date First Named Inventor Goddard, et al. **Art Unit** 1647 Sandra L. Wegert Examiner GNE.3230R1C69 Attorney Docket No.

10/063,596

(Multiple sheets used when necessary) SHEET 1 OF 2

Examiner Initials	Cite No.	Document Number Number - Kind Code (if known) Example: 1,234,567 B1	Publication Date MM-DD-YYYY	Name of Patentee or Applicant	Pages, Columns, Lines Where Relevant Passages or Relevant Figures Appear

FOREIGN PATENT DOCUMENTS						
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Examiner Initials No.		i idam (book magazing jayragi carigi cympacilim catalog atc i date badels) ydillifie-issile i	
	3	CONRADS, et al., "A Combined Proteome and Microarray Investigation of Inorganic Phosphate-induced Pre-osteoblast Cells" <i>Mol. Cell Proteomics</i> , 4(9):1284-1296 (2005).	
	4	CZUPALLA, et al., "Comparative study of proteinand mRNA expression during osteoclastogenesis" Proteomics 5:3868-3875 (2005).	
	5	GINESTIER, et al. 2002. "Distinct and Complementary Information Provided by Use of Tissue and DNA Microarrays in the Study of Breast Tumor Markers" <i>Am. J. Pathol.</i> , 161:1223-1233.	
	6	KAWAMOTO et al., "Expression Profiles of Active Genes in Human and Mouse Livers," <i>Gene</i> , 1996 Sep 26;174(1):151-8.	
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	8	LEDERMAN, et al. 1991. "A single amino acid substitution in a common African allele of the CD4 molecule ablates binding of the monoclonal antibody, OKT4." <i>Molecular Immunology</i> , 28(11):1171-1181.	
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	11	SUGG, et al., "Cytoplasmic staining of <i>erb</i> B-2 but not mRNA levels correlates with differentiation in human thyroid neoplasia" <i>Clinical Endocrinology</i> (1998) 49:629-637.	

			
Examiner Signature	/Sandra Wegert/	Date Considered	01/03/2007

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10/063,596 Application No. INFORMATION DISCLOSURE May 3, 2002 Filing Date Goddard, et al. First Named Inventor STATEMENT BY APPLICANT Art Unit 1647 Examiner Sandra L. Wege (Multiple sheets used when necessary) GNE.3230R1C69 Attorney Docket No. SHEET 2 OF 2

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SLW	12	TOKUNAGA, et al. 2000. Application of quantitative RT-PCR using "TaqMan" technology to evaluate the expression of CK 18 mRNA in various cell lines. J. Exp. Clin. Cancer Res., 19(3):375-381.			
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